

PR 11-JAN-1999 US 09/227881, 07-MAY-1999 US 09/306828 P
THAI D NGUYEN, JON R POLANSKY, PU CHEN, HUA CHEN PC
C12N15/09, A61K31/573, A61K45/00, A61P27/06, C12N1/15, C12N1/19, PC

5225 CCAGTATATATAAACCTCTCTGGAGCTGGGCAAGAGCCAGCAAGG 52/1

Db 181 CCGATATATATAACCTCTCTGAGCTCGGCAATGACCAAG 227

Search completed: January 27, 2006, 09:16:53
Job time : 1 secs

GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: January 27, 2006, 09:20:09 ; Search time 0.001 Seconds

(without alignments)
2393.034 Million cell updates/sec

Title: US-09-227-881-34

Perfect score: 5271
Sequence: 1 acccttgcagcttaccctc.....tcgggcacgagccagcaagg 5271

Scoring table: OLIGO_NUC
Gapop 60.0 , Gapext 60.0

Searched: 1 seqs, 227 residues

Word size : 50

Total number of hits satisfying chosen parameters: 1

Minimum DB seq length: 0

Maximum DB seq length: inf

Post-processing: Listing first 1 summaries

Database : fetch34rnpbm.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Match	Length	DB ID	Description
1	227	4.3	227	1	US-10-244-633-38

ALIGNMENTS

```

RESULT 1
US-10-244-633-38
; Sequence 38, Application US/10244633
; Publication No. US2003068640A1
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; APPLICANT: Chen, Pu
; APPLICANT: Chen, Hua
; TITLE OF INVENTION: Nucleic Acids, Kits, And Methods For The Diagnosis,
; TITLE OF INVENTION: Prognosis And Treatment Of Glioma And Related
; TITLE OF INVENTION: Disorders
; FILE REFERENCE: 07425.0057.US01
; CURRENT APPLICATION NUMBER: US/10/244,633
; CURRENT FILING DATE: 2002-09-17
; PRIOR APPLICATION NUMBER: US/09/306,828
; PRIOR FILING DATE: 1999-05-07
; PRIOR APPLICATION NUMBER: US 09/227,881
; PRIOR FILING DATE: 1999-01-11
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Microsoft Word 97
; SEQ ID NO 38
; LENGTH: 227
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-244-633-38

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Query Match      4.3%; Score 227; DB 1; Length 227;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 227; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 5045 AACTATTATGGGGTATGGGTCATTAATGGATGTTCTTTTAAAGAAAGAACTCCAAA 5104
DB 1 AACTATTATGGGGTATGGGTCATTAATGGATGTTCTTTTAAAGAAAGAACTCCAAA 60
QY 5105 CAGACTTCTGGAAGGTTATTTCTTAAGATCTTGCTGCGACGCGTGAAGGCAACCCCTG 5164
DB 61 CAGACTTCTGGAAGGTTATTTCTTAAGATCTTGCTGCGACGCGTGAAGGCAACCCCTG 120
QY 5165 TGCACAGCCGCCACCCAGCTCAGTGGCCACCTTGCTTCCCATGAAAGGCTGCTC 5224
DB 121 TGCACAGCCGCCACCCAGCTCAGTGGCCACCTTGCTTCCCATGAAAGGCTGCTC 180
QY 5225 CCCAGTATATATAAAGCTCTGAGGCTGGGGGATGAGCCAGCAAGG 5271
DB 181 CCCAGTATATATAAAGCTCTGAGGCTGGGGGATGAGCCAGCAAGG 227

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Search completed: January 27, 2006, 09:20:11
Job time : 1 secs